

Table S1. Major Type Phage (MTPH) reported previously for typing *M. tuberculosis* isolates

MTPH ¹	Phage	Propagation host ¹	Cluster	%GC	Accession #	Reference
1	AG1	<i>M. kansasii</i> At7	N/A	N/A	N/A ²	(1)
2	DS6A	<i>M. tuberculosis</i> H37Rv	Sin	68.40	JN698994	(2)
3	GS4E	<i>M. tuberculosis</i> H37Rv	A2	63.4	AP018479	(3)
4	BK1	<i>M. smegmatis</i> ATCC 607	A1	63.4	AP018477	(3)
5	BG1	<i>M. intracellulare</i> P-17	N/A	N/A	N/A	(4)
6	D34	Uncharacterized <i>M. sp.</i> F130	N/A	N/A	N/A	(5)
7	DNA III	<i>M. tuberculosis</i> H37Rv	G1	67	KC787108.1	(6)
8	X20	<i>M. tuberculosis</i> H37Rv	N/A	N/A	N/A	(7)
9	PH	<i>M. tuberculosis</i> H37Rv	N/A	N/A	N/A	(8)
10	Clark	<i>M. smegmatis</i> ATCC 607	G1	66	KC787112.1	(6)
11	Sedge	<i>M. smegmatis</i> ATCC 607	G1	66	KC787111.1	(6)
12	Legendre	<i>M. smegmatis</i> ATCC 607	G1	67	KC787109.1	(6)

¹MTPH and propagation host are as reported previously (9, 10)

²N/A, Not Available

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